Serval r	Number: 09/726, 643 Edited by: Edited by: Werlfied by: (8TIC st.
PE	Changed the margins in cases where the sequence text was "wrapped: down to the next line. #3
22 2000	Edited a format error in the Current Application Data section, specifically:
& TRAINEMENT	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was   the prior application data; or other
	Added the mandatory heading and subheadings for "Current Application Data".
	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
	Changed the spelling of a mandatory field (the headings of subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included:
	Deleted extra, invalid, headings used by an applicant, specifically:
<u>J</u>	Deleted: ☐ non-ASCII "garbage" at the beginning end of files; ☐ secretary initials/filename at end of file: ☐ page numbers throughout text; ☐ other invalid text, such as
	Inserted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
	Corrected an error in the Number of Sequences field, specifically:
	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
	Deleted ending stop codon in amino acid sequences and adjusted the *(A)Length: field accordingly (error due to a Patentin bug). Sequences corrected:
	Other:

Action. DO NOT send a copy of this form.

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/726,643

DATE: 12/28/2000
TIME: 12:25:16

Input Set : A:\Pto.amc

Output Set: N:\CRF3\12282000\I726643.raw

```
2 <110> APPLICANT: Ruben et al.
      4 <120> TITLE OF INVENTION: 26 Human secreted proteins
      6 <130> FILE REFERENCE: PZ040P1
C--> 8 <140 > CURRENT APPLICATION NUMBER: US/09/726,643
     9 <141> CURRENT FILING DATE: 2000-12-01
    11 <150> PRIOR APPLICATION NUMBER: PCT/US00/15187
    12 <151> PRTOR FILING DATE: 2000-06-02
    14 <150> PRIOR APPLICATION NUMBER: 60/137,725
    15 <151> PRIOR FILING DATE: 1999-06-07
    17 <160> NUMBER OF SEQ ID NOS: 190
    19 <170> SOFTWARE: PatentIn Ver. 2.0
    22 <210> SEQ ID NO: 1
     23 <211> LENGTH: 733
    24 <212> TYPE: DNA
     25 <213> ORGANISM: Homo sapiens
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     29 aattegaggg tgeacegtea gtetteetet teececcaaa acceaaggae accetcatga
                                                                                120
     30 totocoggae tootgaggte acatgogtgg tggtggaegt aagccaegaa gaccetgagg
                                                                                240
     31 tcaagttcaa ctggtacgtg gacggcqtgg aggtgcataa tgccaagaca aagccgcggg
     32 aggageagta caacageaeg taccgtytgg teagegteet caccgteetg caccaggact
                                                                                300
     33 gqctgaatgg caaggagtac aagtgcaagg tetceaacaa agcceteeca acceecateg
                                                                                360
     34 agaaaaccat ctccaaagcc aaagggcagc cccgagaacc acaggtgtac accetycccc
                                                                                480
     35 catceeggga tgagetgace aagaaceagg teageetgae etgeetggte aaaggettet
                                                                                540
     36 atccaaqcqa catcqccqtq qagtqgqaga qcaatggqca gccggagaac aactacaaga
     37 ccacgcetce cgtgetggac tecgacgget cettetteet ctacagcaag etcaccgtgg
                                                                                600
     38 acaagagcag gtggcagcag gggaacgtct totcatgctc cgtgatgcat gaggctctgc
                                                                                720
     39 acaaccacta cacgcagaag ageoteteec tytotooggq taaatgagtg cgacggccgc
                                                                                733
     40 gactetagag gat
     43 <210> SEQ ID NO: 2
     44 <211> LENGTH: 5
     45 <212> TYPE: PRT
     46 <213> ORGANISM: Homo sapiens
     48 <220> FEATURE:
     49 <221> NAME/KEY: Site
     50 <222> LOCATION: (3)
     51 <223> OTHER INFORMATION: Xua equals any of the twenty naturally ocurring L-amino acids
53 <400> SEQUENCE: 2
54 Trp Ser Xaa Trp Ser
     55 1
     57 <210> SEQ ID NO: 3
     58 <211> LENGTH: 86
     59 <212> TYPE: DNA
     60 <213> ORGANISM: Artificial Sequence
W--> 61 <220> FEATURE:
     62 <221> NAME/KEY: Primer_Bind
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63 <223> OTHER INFORMATION: Synthetic sequence with 4 tandem copies of the GAS binding site found in

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```
the 1RF1 promoter (Rothman et al., Immunity 1:457-468 (1994)), 18 nucleotides
              complementary to the SV40 early promoter, and a Xho I restriction site.
     68 grgcctcgag atttccccga aatctagatt tccccgaaat gatttccccg aaatgatttc
     69 occgaaatat etgecatete aattag
     72 <210> SEQ ID NO: 4
     73 <211> LENGTH: 27
     74 <212> TYPE: DNA
     75 <213> ORGANISM: Artificial Sequence
W--> 76 <220> FEATURE:
     77 <221> NAME/KEY: Primer_Bind
     78 <223> OTHER INFORMATION: Synthetic sequence complementary to the SV40 promter; includes a Hind II1
             restriction site.
     81 <400> SEQUENCE: 4
                                                                                27
     82 geggeaaget ttttgeaaag eetagge
     85 <210> SEQ ID NO: 5
     86 <211> LENGTH: 271
     87 <212> TYPE: DNA
     88 <213> ORGANISM: Artificial Sequence
W--> 89 <220> FEATURE:
     90 <221> NAME/KEY: Protein_Bind
     91 <223> OTHER INFORMATION: Synthetic promoter for use in biological assays; includes GAS binding
            sites found in the IRFL promoter (Rothman et al., Immunity 1.457-468 (1994)).
     94 <400> SEQUENCE: 5
                                                                                60
     95 etegagattt eccegaaate tagattteee eqaaatgatt teecegaaat gattteeeeg
     96 aaatatotgo catotoaatt agtoagoaac catagtooog cooctaacto egocoateec
                                                                               120
     97 geocctaact regreeaft eegeceatte teegeceat ggetgaetaa tittittat
                                                                               180
                                                                               240
     98 ttatqcaqaq gccgaqqccq cctcggcctc tgagctattc cagaaqtagt gaggaggctt
     99 ttttggagge ctaggetttt geaaaaaget t
     101 <210> SEQ ID NO: 6
     102 <211> LENGTH: 32
     103 <212> TYPE: DNA
     104 <213> ORGANISM: Artificial Sequence
W--> 105 <220> FEATURE:
     106 <221> NAME/KEY: Primer_Bind
     107 <223> OTHER INFORMATION: Synthetic primer complementary to human genomic EGR-1 promoter sequence
              (Sakamoto et al., Oncogene 6:867-871 (1991)); includes a Xho I restriction site.
     108
     110 <400> SEQUENCE: 6
     111 geget.egagg gatgaeageg atagaaeeee gg
                                                                                 32
     114 <210> SEQ ID NO: 7
     1.15 <211> LENGTH: 31
     116 <212> TYPE: DNA
     117 <213> ORGANISM: Artificial Sequence
W--> 118 <220> FEATURE:
     119 <221> NAME/KEY: Primer_Bind
     120 <223> OTHER INFORMATION: Synthetic primer complementary to human genomic EGR-1 promoter sequence
               (Sakamoto et al., Oncogene 6:867-871 (1991)); includes a Hind III restriction
     121
     122
               site.
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124 <400> SEQUENCE: 7

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Input Set : A:\Pto.amc

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                                                                                  31
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    129 <211> LENGTH: 12
     130 <212> TYPE: DNA
    131 <213> ORGANISM: Homo sapiens
     133 <400> SEQUENCE: 8
     134 ggggaettte ee
                                                                                  12
     137 <210> SEQ 1D NO: 9
     138 <211> LENGTH: 73
     139 <212> TYPE: DNA
     140 <213> ORGANISM: Artificial Sequence
W--> 141 <220> FEATURE:
     142 <221> NAME/KEY: Primer_Bind
     143 <223> OTHER INFORMATION: Synthetic primer with 4 tandem copies of the NF-KB binding site
              (GGGGACTTTCCC), 18 nucleotides complementary to the 5' end of the SV40 early
               promoter sequence, and a XhoI restriction site.
     145
     147 <400> SEQUENCE: 9
     148 geggeetega ggggaettle eeggggaelt teeggggaet tteegggaet tteeateetg
                                                                                  60
     149 ccatctcaat tag
                                                                                  7.3
    152 <210> SEQ ID NO: 10
     153 <211> LENGTH: 256
     154 <212> TYPE: DNA
    155 <213> ORGANISM: Artificial Sequence
W--> 156 <220> FEATURE:
    157 <221> NAME/KEY: Protein_Bind
     158 <223> OTHER INFORMATION: Synthetic promoter for use in biological assays; includes NF-KB binding
              sites.
     161 <400> SEQUENCE: 10
    162 ctcgaggga ctttcccggg gactttccgg ggactttccg ggactttcca tctgccatct
                                                                                 6.0
     163 caattagtea geaaccatag tecegeeest aacteegees atecegees taacteegee
                                                                                 120
     164 cagttoegoe cattetoogo cocatggotg actaatititi titatitatig cagaggooga
                                                                                 180
    165 ggccgcctcg gcctctgagc tattccagaa gtagtgagga ggcttttttg gagqcctagg
                                                                                 240
     166 cttttgcaaa aagett
                                                                                 256
     169 <210> SEQ ID NO: 11
    170 <211> LENGTH: 2318
    171 <212> TYPE: DNA
    172 <213> ORGANISM: Homo sapiens
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     175 cagacetogg acgagagege conggggage toggagegeg tgraegegig geakanggag
                                                                                 60
     176 aaggecagty eccagettga aggitetgic accittigea qiqqiecaaa tqagaaaaaa
                                                                                 120
    177 gtggadaatg ggaggcatga aatacatett ttegttgttg ttetttettt tgetagaagg
    178 aggcaaaaca gagcaagtaa aacattcaga gacatattgc atgtttcaag acaagaagta
                                                                                 240
    179 cagagiggt gagagatgge atectiaect ggaacettat gggttggttt actgegtgaa
                                                                                 300
    180 etgeatetge Leagagaatg ggaatgtget tigeageega gteagatgte caaatgttea
     181 tigoctitot cotgigoala ticotoatot gigotgocot egofgocoag aagaeleett
                                                                                 420
    182 acceeeagtg aacaataagg tgaceageaa gtottgrgag taeaatggga caarttarea
                                                                                 480
    183 acatggagag etgittegtag etgaaggget ettiteagaat eggeaaceea uteaatgeae
                                                                                 540
    184 ccaqtgcagc tgttcgqagq gaaacgtqta ttgtggtctc aagacttqcc ccaaattaac
                                                                                 600
    185 etgtgeette coagtetetg ttocagatte etgetgeegg gtatgeagag gagatggaga
                                                                                 660
```

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			at aut and at	ottooggoaa	oetagosaes	garnaggaag	720
186	actigueatgq	gaacattetg	arggradacar	tocaccaage	cetgecaata	gagaagcaag	780
187	acattettae	cacegetete	actatgatee	tocaccaage	cgacaggerg	anggeogee	840
188	cogetttect	ggggccagaa	greaecqqqq	agetettatg	gatteecage	etatttaana	900
189	aaccattgtg	caaattgtca	теаатаасаа	acacaagcat	ggacaagege	ttaaaattat	960
190	tggaaagacc	tattctcatg	gegagteetg	gcacccaaac	between	teggeatige	1020
191	ggagtgtgtg	ctatgtactt	gtaatgtcac	caagcaagag	Egraagaaaa	teeactycec	1.020
192	caatcqatac	ccctgcaagt	atceteaaaa	aatagacgga	aaatgetgea	aggrated	
193	agaagaactt	ccaggccaaa	getttgacaa	taaaggetae	ttetgegggg	aagaaacgat	1140
194	gcctgtgtat	gagtetgtat	tcatggagga	tggggagaca	accagaaaaa	tagcactgga	1200
195	gactgagaga	ccacctcagg	tagaggtcca	cgtttggact	attegadagg	geattetea	1260
1.96	gcacttccat	atigagaaga	tetecaagag	gatgtttgag	gagetteete	acttcaaget	1320
197	ggtgaccaga	acaaccctga	gecagtggaa	gatetteace	gaaqgaqaaq	ctcagatcag	1380
198	ccagatgtgt	teaagtegtg	tatgeagaac	agagettgaa	gatttagtca	aggttttgta	1.440
199	cctggagaga	tctgaaaagg	gccactgtta	ggcaagacag	acagtattyg	ataggqtaaa	1500
200	qcaaqaaaac	teaagetgea	getggaetge	aggettattt	tgctLaagtc	aacagtgccc	1560
201	taaaactcca	aactcaaatg	cagtcaatta	ttcacqccat	gcacagcata	atttgctcct	1.620
202	ttatatatat	gratatatat	gtgtgtgtqt	gtgtggtaaa	qqqqqgaaqq	tyttatgegg	1680
203	ctgetcecke	cqt.cccaqaq	qt.qqcaqt.qa	Ltccataatg	tggagactag	Laactagato	1740
204	chaaggcaaa	gaggtgtttc	teettetgga	tgattcatcc	caaagcette	ccacccaggt	1800
205	attetetaaa	agettageet	taaqaqaaca	cqcagagagt	thocotagat	atactectge	1860
206	checaggiac	tgugacacac	etttgcaaaa	tgctgtggga	agcaggaget	ggggagetyt	1920
207	of taagleaa	agtagaaacc	ctccagtgtt	LggLgtLgLg	Lagagaatag	gacataggqt	1980
203	азапапписса	agetgeetgt	agt tautaga	gaagaatgga	taraattett	cttgtgtatt	2040
200	tatttatate	ataaacactt	ggaacaacaa	agaccataag	catcatttag	cagttgtage	21.00
210	cattttttaa	traactcato	taaacaagta	agagtaacat	aacaqtatta	ccetttcact	2160
210	attatasasa	gacatgtacc	taattalggt	acttatttat	gragt.cact.g	Latitictoga	2220
21.1	tettetaaatt	ant annual	ttaatittaa	aaaatcaaaa	аалаалааааа	aaaagt cgac	2280
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	<210> SEQ		gragiagrag	cagcagga			
	<211> LENG						
	<21.2> TYPE		anniana				
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223	atagetecat	ecagectgag	aaacaagccg	ggtggcLgag	ccaggoogra	gagatotoot	180
224	tgacgggccc	aacagaccca	Egnigearee	agagacet.cc	et et accese	ggesteeteet	240
	agetgtgete	ctadeceree	Liggeacege	ctggggagag	grig riggionale	teatutusat	300
226	ggagcagget	eegatggeeg	gagnectgaa	caggaaggag	agrittettge	recterent	360
227	geacaaccge	ctgcgcaget	gagtecaace	acat.gaggat	gacatigogga	ggetggaetg	420
228	gagtgacage	etggeeeaac	tggctcaage	cagggcagcc	cucuguggaa	ucecaacce	
229	gagectggcg	teeggeetgt.	ggognacect	gcaagtgggc	Lggaacat.gc	agelgetgee	480
230	cgegggettg	gegteetttg	ttgaagtggt	cagectutgg	thighagagg	ggcagcggta	540
231	cagecaegeg	gcaggagagt	gtgetegeaa	agacacatge	acchaettaca	egeageregu	600
232	gtgggddaid	teaageeage	Laggetatag	geggeacetg	tgctctgcag	gccaggcagc	660
233	gatagaagee	lttgtclgtg	ectactocce	cggaggcaac	Lgggaggtca	acgggaagac	720
234	aatcatcccc	tataagaagg	gtgcctggtg	ttegetetge	acagecagtq	totoaggetg	780
235	intheaaagre	taggaccata	caggggggct	intgtgaggtic	-cccaggaatc	cttgtcgcat	840
236	gagetgecag	aaccatqgac	greteaacat	cagcacctgc	cactgccact	gthacaatgg	900
237	etacacggge	agatactgcc	aagtgaggtg	cageotycay	tylylycacy	geeggtteeg	960

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238	qqaqqaqaq	tactcataca	tetgtgacat	cggctacggg	ggagcccagt	gtgecaccaa	1020
239	ggtgcatttt	cccttccaca	cclgtgacct	gaggategae	ggagactgct	teatggtgte	1.080
240	t t.cagaggca	gacacctatt	acagagecag	gatgaaatgt.	cagaggaaag	gcggggtgct	1.140
241	ggcccagate	aagagccaga	aagtgcagga	catectegee	ttctatctgg	gccgcctgga	1.200
242	gaccaccaac	gaggtgattg	acaglgactt	cgagaccagg	aacttetgga	togggetcac	1260
243	chacaagacc	gccaaggact	ccttccyctg	ggccacaggg	gagcaccagg	ccttcaccag	1320
244	tittacetti	gggcagcctg	acaaccacqq	gtttggcaac	tgcgtggage	tgcaggcttc	1.380
245	agetgeette	aactggaaca	accagegetg	caaaacccga	aaccgttaca	totgocagtt.	1.440
246	tacccaggag	cacatetece	ggtggggccc	agggteetga	ggcctgacca	catggetece	1500
247	tracetacee	Laggageace	ggctctgctt	acctiftccgc	ccacctgtct	ggaacaaggg	1560
248	ccaggitaag	accacatgcc	teatgtecaa	agaggtetea	gacettgeae	aatgccagaa	1620
249	attaggcaga	gagaggeagg	gaggeeagtg	agggccaggg	agtgagtgtt	agaagaaget	1680
250	gagagagatta	acctactitt	gattgggaag	atgggettea	attagatggc	gaaggagagg	1.740
250	acaccaccac	tootccaaaa	aggetyetet	cttccacctq	geneagaece	tqt.qqqqcag	1800
252	canaact tee	clatageata	aaccccacag	ggtattaaat	tatqaatcaq	ctgaaaaaaa	1860
252	заааааааааа	ааааааааааа	аааааааааа	aaaaaaaaaa	aaaaaaaaa	aaaaaaaaa	1.920
	aaa						1923
	<210> SEQ :	m No. 13					
	<211> LENG'						
	<212> TYPE						
	<213> ORGAI		sapiens				
	<400> SEQUI		,				
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264	tetatteeet	cctuagectu	agececttae	ettectgace	ccatgaagca	cacactgget	1.20
265	chachagete	ccctactaaa	cetgageetg	aggetugece	tgagtcagct	ggetgeagag	180
266	accacagaet	genagticet	Lggcccggca	gagcacctga	cat Leaccec	agcagecagg	240
267	acconstance	taacccctca	agttegtgeg	ccaggactice	Lagactecet	ctatggcacc	300
268	atacaccact	tectetedat	ggtqcaqctc	aatcetttee	effeagagtt	ggtaaaggcc	360
260	ctactgaatg	agetqueete	cataaaaata	aatgaggtgg	tgeggtaega	qqeqqqctac	420
270	ataatataa	ctataateac	gggcetetae	ctactactaa	Lacceactac	egagetttye	480
270	trotactact	accact acca	acadedetac	gggggacgag	t.gaagacaga	gcacaaggcg	540
27.1	ctagectata	ageageageaa	ceteatagte	tteetgetge	tgaccaccet	cttqctqctq	600
272	attaatataa	tetatacett	tatcaccaac	caucucacuc	atuaacagat	gggccccagc	660
273 274	ategaracea	tacctuagec	cetacteage	ctctggggcc	Lagitetetga	figicocceaa	720
275	dadotacada	centageaca	gcaattetee	ctgcccagg	agcaagtete	agaggagetg	780
275	gageegeagg	atataaacat	taggaucaca	atccacactc	ageteaggag	ctccgtgtac	840
277	gacagegrea	gagagagag	cadtttaggc	caggirectige	aggleteegt	gcaccacctg	900
277	eaching tra	atgetac it	ontananclu	cadaccadac	ageaggaect.	ggagecagec	960
270	ntacoddaac	acqueaccq	- cetect Equi	ctactacaga	aggecaggtg	ccagagagat	1020
277	tatacaunga	ecctoageta	- dacecacaca	ctggagetgg	gtactgactt	cagecaggtg	1080
200	contatataa	accalatect	ggaccadela	aaaggtgtcc	ecgaggecaa	ettetecage	1140
201	atracticeagg	addadaacad	- cacetteaac	accettecag	ccetagetae	catgoagaca	1200
202	toggoodigg	- tacaadadat	- dedication -	ataacccaac	adccggaagg	qqtqaqaca	1260
203	- cccaycycyg	antipecaea	- cit doadoca	act Leceart	aducceador	actgcaggag	1320
204	- clygelyddy - alagaegae	991.0000999	- ctacctacea	daggtucada	galacdagac	ctacaggt.gg	1380
285	giggaggaga	geageeyeee	- ctachigada	- ctattenton	tactctacaa	cotqutqqqc	1440
285	aloguagaci	graugerard acatotadad	- catatataa	adddaedaec	- сеадесяесе	agaagecaag	1500
287	negairergg	geacetyggg	- codycocyco	- agggacgacc	tragetteet	ctitgctgca	1560
288	ddedadderd	- gagedegett	- concorregged	- gg cy cygyddd - at aat aast a	- acagent des	gacgctggtg	1620
289	econtrated	udduggugut	e, good octio	cuggugggug	geaucycycu	9409009909	1.021



## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/726,643 TIME

DATE: 12/28/2000 TIME: 12:25:17

Input Set : A:\Pto.amc

Output Set: N:\CRF3\12282000\I726643.raw

L:8 M:270 C: Current Application Number differs, Replaced Current Application Number L:54 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 L:61 M:283 W: Missing Blank Line separator, <220> field identifier L:76 M:283 W: Missing Blank Line separator, <220> field identifier L:89 M:283 W: Missing Blank Line separator, <220> field identifier L:105 M:283 W: Missing Blank Line separator, <220> field identifier L:118 M:283 W: Missing Blank Line separator, <220> field identifier L:141 M:283 W: Missing Blank Line separator, <220> field identifier  $L:156\ \dot{M}:283\ W:$  Missing Blank Line separator, <220> field identifier L:586 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 L:668 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 L:673 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 L:897 M:341 W: (46) "n" or "Xaa" used, for SEQ 1D#:25 L:898 M:341 W: (46) "n" or "Xaa" used, for SEQ 1D#:25 L:1052 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 L:1154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 L:1162 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 L:1237 M:341 W: (46) "n" or "Xaa" used, for SEO 1D#:33 L:1301 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 L:1302 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 L:1303 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 L:1379 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 L:1547 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 L:1551 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:39 L:2197 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49 L:2529 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53 L:2625 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:54 1:2993 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60 L:2996 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60 L: 388 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69 L:3391 M:341 W: (46) "n" or "Xaa" used, for SEQ 1D#:69 L:3394 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69 L:3481 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:70 L:3863 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81 L:3884 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:82 L:4812 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:107 L:4833 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:107 L:4921 M:311 W: (46) "n" or "Xaa" used, for SEQ ID#:112 L:4964 N:341 W: (46) "n" or "Xaa" used, for SEQ ID#:114 L:4979 M:341 W: (46) "n" or "Xaa" used, for SEQ 1D#:115 L:5857 M:311 W: (46) "n" or "Xaa" used, for SEQ 1D#:151 L:5926 H:341 W: (46) "n" or "Xaa" used, for SEQ ID#:155 L:5958 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:156 L:5961 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:156 L:6006 M:311 W: (46) "n" or "Xaa" used, for SEQ 1D#:158 1.:6009 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:158